³**Floristic composition in chronosequence in** ⁴**Atlantic Rainforest fragments**

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10 : 11 **ABSTRACT**

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> Secondary forests play an important role in tropical landscapes and have important ecological functions such as the ability to accumulate biomass. Although the literature points to the convergence between primary and secondary forests, however there are few studies in Atlantic Rainforest in a chronosequence to show it. This study aimed to characterize the changes of floristic composition in a chronosequence (5, 16, 24, 30 years of regeneration and mature forest) in the Atlantic Rainforest. He sought to answer the following questions: Young forests tend to converge resembling floristically with mature forest? In case of this floristic convergence from what age these young forests present rate of growth closer forms of mature forest? The fact that about 35% of the tree species are present in young and mature forests suggests that the floristic composition of young forests tends to resemble the mature forest. The results suggest that from 16 young forests tended to converge with the mature forest in terms of the proportion of growth forms.

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14 *Keywords: Floristic convergence, Young forests, Growth forms, Secondary rainforest.*

16 **1. INTRODUCTION**

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18 Secondary forests play an important role in tropical landscapes. These ecosystems are 19 sources of timber and non-timber products and generally perform important ecological 20 functions, such as the ability to accumulate biomass and nutrients at high rates, maintain
21 biogeochemical cycles and water conservation at levels comparable to primary forests [1.2]. 21 biogeochemical cycles and water conservation at levels comparable to primary forests [1,2].
22 Current research on succession in secondary forests in the tropical regions occurs especially Current research on succession in secondary forests in the tropical regions occurs especially 23 in studies in chronosequences [3].

24 Regarding succession [4] proposed that it is a highly ordered and predictable process, 25 emphasizing the sequential substitution of herbaceous species by shrubs and finally 26 arboreal species. Differently, [5] suggested that the sequence does not have this fixed and 27 regular sequence, being influenced by the composition of the community, which is 28 determined by stochastic factors.

29 In relation to floristic composition throughout the succession, [6] described two alternative 30 scenarios. The floristic rotation, with a broadly clementsian view, in which each group of 31 species colonizes the place at certain stage of development, thus making conditions 32 unsuitable for itself and more appropriate for colonization of the next group, at the final stage 33 of succession convergence with the region's mature vegetation. Other scenario is the 34 hypothesis of initial floristic composition, which proposes that after the abandonment of a 35 site, development unfolds from the initial flora; [6] concluded that the secondary succession

36 is determined more by the initial floristic composition of an area than by the floristic rotation 37 proposed by [4].

38 At present, it is known that natural landscapes, in general, have a historical complex of land 39 use disturbances [7]. These authors argued that succession represents a series of 40 unpredictable events that results from interactions between individuals and the abiotic 41 environment, involving different patterns (convergence or divergence), mechanisms and 42 causes that lead to complex paths that will determine the processes of recovery of the local 43 vegetation. The trajectory followed by the sequence depends on the initial conditions and 44 stochastic events. The results of chronosequences found in tropical forests by several 45 authors [2,8] have reported floristic convergence with mature vegetation.

46 Although the literature cites some papers that point to convergence, it is known that there 47 are studies reporting the floristic divergence along the chronosequence [9,10]. The authors 48 argued that this may be due to less similarity of habitats and/or landscape effects [11]. On 49 the landscape effect, [12] found that in landscapes with less than 10-30% of vegetation 50 covering, there are negative effects on the persistence of the species due to the combined 51 effect of landscape area and geometry, which may lead to convergent or divergent 52 trajectories, according to the degree of vegetative cover.

53 From the perspective of the current literature, there is still a great deal of research to be 54 done. The present research aims to characterize the changes in floristic composition in a 55 chronosequence (5, 16, 24, 30 years of regeneration and mature forest). Thus, we sought to 56 answer the following questions: From what age do young forests present a proportion of 57 growth forms closer to the mature forest? Do young forests tend to diverge or converge 58 floristically with mature forest?

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60 **2. MATERIAL AND METHODS**

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62 **2.1. Local study and selection of areas**

63 The fragments of mature and young forest studied are located at the São José Plant (USJ) 64 (07° 54' 17'' - 07° 54' 41''S, 34° 54' 17 "- 35° 05' 07''), located in the Zona da Mata Norte de 65 Pernambuco, in the domain of the Dense Ombrophylous Forest [13]. The property occupies 66 270 km², where it has forest remnants that cover 24% of the landscape, being the average 67 size of the fragments of 61 ha, occupying the less useful areas for planting [14].

68 The local climate is As' (hot and humid), with an annual average temperature of 24.9 °C, 69 average rainfall of 1687 mm (Laboratory of Meteorology of Pernambuco - LAMEPE/ITEP).

70 Young forests with 5, 16, 24 and 30 years of regeneration were selected from aerial
71 photographs of the 1960s, 1970s, 1980s, 2005 satellite images and interviews with former photographs of the 1960s, 1970s, 1980s, 2005 satellite images and interviews with former 72 residents. As a comparison, an area of mature forest (over 60 years old) studied by [15] was 73 selected. It is important to point out that mature forest is the one in which we can guarantee, 74 based on cartographic material and interviews, that there was no intervention in the last 60 75 years.

76 **2.2. Data collect**

77 Thirty plots of 10×10 m were sampled in each forest of the chronosequence, as well as the 78 floristic survey near the plots. The collection was carried out by the walking method [16], 79 according to the usual collection techniques for plants [17]. The studied habits were: 1) 80 arboreal; 2) shrub; 3) grass; 4) bindweed (woody) and 5) epiphyte.

81 **2.3. Analysis and processing of data**

82 The species sampled were identified with the help of experts and comparisons with 83 collections deposited at the Herbarium Sergio Tavares (HST) and Dárdano de Andrade Lima 84 (IPA). The exsiccates were placed in the Herbarium Professor Vasconcelos Sobrinho 85 (PEUFR). The classification of angiosperm families followed the recommendations of [18].

86 To test the possible convergence or divergence of habits between the different dropout ages 87 and the proportion between the different habits during the chronosequence, one-way 88 ANOVA was applied, followed by the post-hoc Tukey test through the PAST 2.01 software 89 [19].

90 In order to verify if there is floristic convergence between the young forests and the mature 91 forest the degree of floristic similarity between the tree species in the chronosequence was 92 evaluated. To do this, we performed: 1) a clustering analysis using the Jaccard similarity 93 index and the mean linkage method per group (UPGMA) [20]; 2) Principal component 94 analysis (PCA), Multidimensional linear analysis technique [21], and 3) non-metric 95 multidimensional scaling (NMDS) [22]. Based on the tree species present in the five ages, a 96 presence/absence matrix was elaborated to analyze if there would be indicator species for 97 each age, using TWINSPAN (Two Way Indicator Species Analysis) analysis [23] using PC-98 ORD version 4.0 [24]. The option to analyze only trees occurred because it is the best 99 registered habit in the five areas.

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101 **3. RESULTS AND DISCUSSION**

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103 In the five fragments sampled, 198 taxa were recorded, distributed by 131 genera and 59 104 families (Table 1). In the area of 5 years, 66 species and 42 families occurred, being 105 Myrtaceae and Melastomataceae the most representative. In the forest of 16 years of 106 abandonment, we found 75 species and 44 families, with greater representation of 107 Fabaceae, especially the subfamilies Papilionoideae and Mimosoideae. In the area of 24 108 years, 88 species and 51 families occurred, Myrtaceae being the one with the greatest 109 number of species. In the 30 years, with 75 species and 41 families, with highlight to 110 Myrtaceae and Fabaceae (Mimosoideae) families, due to the greater species richness. In 111 the mature forest, there were 95 species and 52 families, with Myrtaceae, Sapindaceae and 112 Rubiaceae with the largest number of species (Table 1).

113 The trends observed during the chronosequence in the fragments at the São José Plant 114 reveal that in the youngest forest (5 years) there were several herbs with a short life cycle 115 and an increase in the number of tree species and decrease of the herbs in the more 116 advanced stages (16, 24, 30 and mature), what was pattern observed in other 117 chronosequence studies [25,26,27].

118 Tree habit showed the highest number of families (43) and species (124), from these 124 119 species, 51 occurred in the young forests and 72 in the mature forest, and the two groups 120 shared 43 species. The families that presented the greatest wealth of trees were Myrtaceae 121 and Fabaceae (Mimosoideae), with 15 and 10 species, respectively. There was an increase 122 in the number of species in the families Sapindaceae, Annonaceae, Myrtaceae and 123 Fabaceae throughout the chronosequence; from these 124 tree species recorded in young 124 forests and mature forest, 15 species (12.9%) were common at all ages (Table 1).

125 The shrub habitat presented 13 families and 39 species. In the herbs group, 12 families and 126 19 species were recorded (Table 1). Among the bindweed, nine families and 16 species 127 were recorded (Table 1). The epiphyte of the hemiparasite type *Psittacanthus dichrous* 128 (Loranthaceae) was found only in the young forest of 5 years.

129 It can be noticed an advance to a more typical pattern of mature forests from 16 years after 130 the abandonment, due to the increase of the trees and decrease of the herbaceous ones, 131 showing the directional substitution of forms of growth.

132 The small trees that occurred in all the young forests are typical species of edges and
133 clearings of forests [28,29], being present in the young forests in sunny areas [30]. 133 clearings of forests [28,29], being present in the young forests in sunny areas [30].

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Table 1. Species recorded in four fragments of young forests (5, 16, 24 and 30) and a mature forest. ARO - Arboreal, TRE - Bindweed, ARB - Bush, ERV - Herb, EPI -**Epiphyte. Young forests (FJ5 = 5 years = 16 years FJ16, FJ24 = 24, FJ30 = 30 years) and mature forest (FM). ^{*} arborescent plant with similar size to tree. Through the analysis of indicator species TWINSPAN: ♦ mature forest; • young forests.**

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141 Significant differences were observed in terms of number of species in tree and herbaceous 142 habits among the different ages studied (Fig. 1). There was a trend of increasing tree 143 species and decreasing of herbs along the chronosequence, where mature forest had the 144 highest number of species $(p \le .02)$ in relation to the young (5 years) . In the 145 chronosequence, another trend was observed, in which the richness of the herbaceous 146 growth form decreased towards the mature forest, where the forest of the initial stage (5 147 years) presented the highest proportion, differing from the other forests studied ($p < .01$) 148 (Fig. 1). In this way, it was observed that from 16 years after the abandonment the 149 proportion of the growth forms were closer to the mature forest.

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159 **Fig. 1. Average number of species per plot with different growth habits in four young** 160 **forests (5, 16, 24 and 30) and a mature forest.** *Letters indicate that the average within each* 161 *habit do not differ by Tukey test (P <0.05) and the same letter are not statistically different from each* 162 *other. Young forests (FJ5 = 5 years = 16 years FJ16, FJ24 = 24, FJ30 = 30 years) and mature forest* 163 *(FM).* 164

165 However, the analysis of TWINSPAN (Table 1) identified in division 1 (eigenvalue 0.911) the 166 mature forest area of the young forests. *Albizia polycephala, Artocarpus heterophyllus,* 167 *Xylopia frutescens, Apeiba tibourbou* and *Vismia guianensis* were arboreal species 168 indicative of young forests, as well as the *Acrocomia sclerocarpa* palm, which was also 169 common in young forests. In the mature forest, only species with more advanced stages 170 were found, such as *Pogonophora schomburgkiana*, *Simarouba amara*, *Guatteria* 171 *schomburgkiana, Maytenus distichophyla, Dialium guianense* and *Lecythis pisonis* (Table 1). 172 The species considered as indicators of mature forest are in agreement with several authors 173 [28,29], since they are species with similar habitat characteristics, being common in the inner 174 of forests and shaded areas.

175 NMDS analysis applied to the tree species in the chronosequence revealed the formation of 176 two groups, isolating the mature forest (FM) (Fig. 2A). The grouping analysis individualized 177 the mature forest (FM) and two groups, one formed by the young forests of 5 (FJ 5) and 16 178 years (FJ 16) of regeneration, with 46% similarity and a second formed by the young forests 179 of 24 (FJ 24) and 30 years (FJ 30), with 50% similarity (Fig. 2B). The two groups composed 180 of young forests (FJ 5 - FJ 16 and FJ 24 - FJ 30) differed from mature forest in floristic
181 composition with 28% similarity. This same pattern was also evident in the principal composition with 28% similarity. This same pattern was also evident in the principal

182 component analyzes (PCA) (Fig. 2C).

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Fig. 2. Multivariate analysis based on the presence and absence of tree species sampled in young and mature forest forests. Ordination by the not metric multidimensional scaling method (NMDS) (A), cluster analysis generated by the Jaccard similarity indices and average link method group (B) and principal component analysis (C). Young forests (FJ_5 = 5 years FJ_16 = 16, FJ_24 = 24, FJ_30 = 30 years) and mature forest (FM).

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191 The results of all multivariate analyzes indicate the formation of two groups of young forests 192 (5 and 16 years old and another 24 and 30 years old), individualizing the mature forests. The 193 literature also points out that the recovery of floristic young forests occurs slowly [31,2]. 194 Regarding this slowness, the fact that young forests and mature forest share 43 tree 195 species, 15 in all forests, suggests that the floristic composition of young forests tends to 196 converge with mature forest, as well as recorded in several researches in tropical forests 197 [2,32,33,34,30].

198 It can be hypothesized that the floristic convergence and the strong influence of the initial 199 floristic composition on the chronosequence flora in the fragments may be related to the fact 200 that this area presents more than 24% vegetation cover, since, according to [12], landscapes 201 above the threshold of 10-30% of covering tend to have positive effects on the persistence of 202 species, leading to different trajectories according to the size and connectivity of the 203 patches. 204

205 **4. CONCLUSIONS**

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207 In the chronosequence studied at the São José Plant, it can be seen that from 16 years, 208 young forests tended to converge with mature forest in proportion of tree and herbaceous 209 species. The distribution of 43 tree species among young and mature forests suggests that 210 the floristic composition of young forests tends to resemble that of mature forest, confirming 211 the hypotheses of floristic convergence.

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213 Floristic convergence may have been influenced by the São José Plant landscape, which 214 has a 24% vegetation cover, facilitating the persistence of the species in the studied 215 fragments. In addition to sharing 43 species, there were 15 species that were present at all 216 ages, supporting the hypothesis of initial floristic composition, in which the species that 217 participate in the succession over time are established from the beginning in the abandoned 218 area. 219

- 220 **COMPETING INTERESTS**
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222 Authors have declared that no competing interests exist. 223

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